



-1-

SEQUENCE LISTING

<110> Thomas, Stephen G
Hedden, Peter
Phillips, Andrew L

<120> Gibberellin 2-Oxidase

<130> 0623.0970000

<140> To Be Assigned

<141> Herewith

<150> PCT/GB99/01857

<151> 1999-06-11

<150> GB 9812821.8

<151> 1998-06-12

<150> GB 9815404.0

<151> 1998-07-15

<160> 16

<170> PatentIn Ver. 2.1

<210> 1

<211> 1318

<212> DNA

<213> Phaseolus coccineus

<400> 1

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<210> 2

<211> 331

<212> PRT

<213> Phaseolus coccineus

<400> 2

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 Thr His Pro Asp Ala Lys Asn Leu Ile Val Asn Ala Cys Arg Asp Phe
 35 40 45
 Gly Phe Phe Lys Leu Val Asn His Gly Val Pro Leu Glu Leu Met Ala
 50 55 60
 Asn Leu Glu Asn Glu Ala Leu Arg Phe Phe Lys Lys Ser Gln Ser Glu
 65 70 75 80
 Lys Asp Arg Ala Gly Pro Pro Asp Pro Phe Gly Tyr Gly Ser Lys Arg
 85 90 95
 Ile Gly Pro Asn Gly Asp Val Gly Trp Val Glu Tyr Leu Leu Leu Asn
 100 105 110
 Thr Asn Pro Asp Val Ile Ser Pro Lys Ser Leu Cys Ile Phe Arg Glu
 115 120 125
 Asn Pro His His Phe Arg Ala Val Val Glu Asn Tyr Ile Thr Ala Val
 130 135 140
 Lys Asn Met Cys Tyr Ala Val Leu Glu Leu Met Ala Glu Gly Leu Gly
 145 150 155 160
 Ile Arg Gln Arg Asn Thr Leu Ser Arg Leu Leu Lys Asp Glu Lys Ser
 165 170 175
 Asp Ser Cys Phe Arg Leu Asn His Tyr Pro Pro Cys Pro Glu Val Gln
 180 185 190
 Ala Leu Asn Arg Asn Leu Val Gly Phe Gly Glu His Thr Asp Pro Gln
 195 200 205
 Ile Ile Ser Val Leu Arg Ser Asn Ser Thr Ser Gly Leu Gln Ile Cys
 210 215 220
 Leu Thr Asp Gly Thr Trp Val Ser Val Pro Pro Asp Gln Thr Ser Phe
 225 230 235 240
 Phe Ile Asn Val Gly Asp Ala Leu Gln Val Met Thr Asn Gly Arg Phe
 245 250 255
 Lys Ser Val Lys His Arg Val Leu Ala Asp Thr Thr Lys Ser Arg Leu
 260 265 270
 Ser Met Ile Tyr Phe Gly Gly Pro Ala Leu Ser Glu Asn Ile Ala Pro
 275 280 285
 Leu Pro Ser Val Met Leu Lys Gly Glu Glu Cys Leu Tyr Lys Glu Phe
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 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Probe

<400> 3

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<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Probe

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gccatgcctt gtccctgagc aagatgattg gctttacaaa gaattcactt ggtctcaata 180
caaatcttct gcttacaag                                     199
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<210> 5

<211> 1318

<212> DNA

<213> Arabidopsis thaliana

<220>

<221> misc_feature

<222> (1243, 1265)

<223> unidentified residue

<400> 5

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atccaaacat gccctcgtga aagcatgcga agacttcggc ttcttcaagg tgatcaacca 180
tggtcggttcc gcagagctag tctctgtttt agaacacgag accgtcgatt tcttctcggt 240
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tggtgctaggg atcaaaccca ggaacacact tagcaagctt gtgtctgacc aaaacacgga 540
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tcccgatcac acttcttct tcttcaacgt tggtgactct ctccagggtg tgacaaatgg 780
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<210> 6

<211> 329

<212> PRT

<213> Arabidopsis thaliana

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35 40 45
Val Ser Ala Glu Leu Val Ser Val Leu Glu His Glu Thr Val Asp Phe
50 55 60
Phe Ser Leu Pro Lys Ser Glu Lys Thr Gln Val Ala Gly Tyr Pro Phe
65 70 75 80
Gly Tyr Gly Asn Ser Lys Ile Gly Arg Asn Gly Asp Val Gly Trp Val
85 90 95
Glu Tyr Leu Leu Met Asn Ala Asn His Asp Ser Gly Ser Gly Pro Leu
100 105 110
Phe Pro Ser Leu Leu Lys Ser Pro Gly Thr Phe Arg Asn Ala Leu Glu
115 120 125
Glu Tyr Thr Thr Ser Val Arg Lys Met Thr Phe Asp Val Leu Glu Lys
130 135 140
Ile Thr Asp Gly Leu Gly Ile Lys Pro Arg Asn Thr Leu Ser Lys Leu
145 150 155 160
Val Ser Asp Gln Asn Thr Asp Ser Ile Leu Arg Leu Asn His Tyr Pro
165 170 175
Pro Cys Pro Leu Ser Asn Lys Lys Thr Asn Gly Gly Lys Asn Val Ile
180 185 190
Gly Phe Gly Glu His Thr Asp Pro Gln Ile Ile Ser Val Leu Arg Ser
195 200 205
Asn Asn Thr Ser Gly Leu Gln Ile Asn Leu Asn Asp Gly Ser Trp Ile
210 215 220
Ser Val Pro Pro Asp His Thr Ser Phe Phe Phe Asn Val Gly Asp Ser
225 230 235 240
Leu Gln Val Met Thr Asn Gly Arg Phe Lys Ser Val Arg His Arg Val
245 250 255
Leu Ala Asn Cys Lys Lys Ser Arg Val Ser Met Ile Tyr Phe Ala Gly
260 265 270
Pro Ser Leu Thr Gln Arg Ile Ala Pro Leu Thr Cys Leu Ile Asp Asn
275 280 285
Glu Asp Glu Arg Leu Tyr Glu Glu Phe Thr Trp Ser Glu Tyr Lys Asn
290 295 300
Ser Thr Tyr Asn Ser Arg Leu Ser Asp Asn Arg Leu Gln Gln Phe Glu
305 310 315 320
Arg Lys Thr Ile Lys Asn Leu Leu Asn
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<210> 7
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 <212> DNA
 <213> Arabidopsis thaliana

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<210> 8
 <211> 341
 <212> PRT
 <213> Arabidopsis thaliana

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 35 40 45
 Cys Glu Glu Phe Gly Phe Phe Lys Val Val Asn His Gly Val Arg Pro
 50 55 60
 Glu Leu Met Thr Arg Leu Glu Gln Glu Ala Ile Gly Phe Phe Gly Leu
 65 70 75 80
 Pro Gln Ser Leu Lys Asn Arg Ala Gly Pro Pro Glu Pro Tyr Gly Tyr
 85 90 95
 Gly Asn Lys Arg Ile Gly Pro Asn Gly Asp Val Gly Trp Ile Glu Tyr
 100 105 110
 Leu Leu Leu Asn Ala Asn Pro Gln Leu Ser Ser Pro Lys Thr Ser Ala
 115 120 125
 Val Phe Arg Gln Thr Pro Gln Ile Phe Arg Glu Ser Val Glu Glu Tyr
 130 135 140
 Met Lys Glu Ile Lys Glu Val Ser Tyr Lys Val Leu Glu Met Val Ala
 145 150 155 160
 Glu Glu Leu Gly Ile Glu Pro Arg Asp Thr Leu Ser Lys Met Leu Arg
 165 170 175

Asp Glu Lys Ser Asp Ser Cys Leu Arg Leu Asn His Tyr Pro Ala Ala
180 185 190

Glu Glu Glu Ala Glu Lys Met Val Lys Val Gly Phe Gly Glu His Thr
195 200 205

Asp Pro Gln Ile Ile Ser Val Leu Arg Ser Asn Asn Thr Ala Gly Leu
210 215 220

Gln Ile Cys Val Lys Asp Gly Ser Trp Val Ala Val Pro Pro Asp His
225 230 235 240

Ser Ser Phe Phe Ile Asn Val Gly Asp Ala Leu Gln Val Met Thr Asn
245 250 255

Gly Arg Phe Lys Ser Val Lys His Arg Val Leu Ala Asp Thr Arg Arg
260 265 270

Ser Arg Ile Ser Met Ile Tyr Phe Gly Gly Pro Pro Leu Ser Gln Lys
275 280 285

Ile Ala Pro Leu Pro Cys Leu Val Pro Glu Gln Asp Asp Trp Leu Tyr
290 295 300

Lys Glu Phe Thr Trp Ser Gln Tyr Lys Ser Ser Ala Tyr Lys Ser Lys
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His Lys Thr Leu Val
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<210> 9
<211> 1008
<212> DNA
<213> Arabidopsis thaliana

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<210> 10
<211> 335
<212> PRT
<213> Arabidopsis thaliana

<400> 10
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      35                      40                      45
Gly Phe Phe Lys Val Ile Asn His Gly Val Arg Pro Asp Leu Leu Thr
      50                      55                      60
Gln Leu Glu Gln Glu Ala Ile Asn Phe Phe Ala Leu His His Ser Leu
      65                      70                      75                      80
Lys Asp Lys Ala Gly Pro Pro Asp Pro Phe Gly Tyr Gly Thr Lys Arg
      85                      90                      95
Ile Gly Pro Asn Gly Asp Leu Gly Trp Leu Glu Tyr Ile Leu Leu Asn
      100                     105                     110
Ala Asn Leu Cys Leu Glu Ser His Lys Thr Thr Ala Ile Phe Arg His
      115                     120                     125
Thr Pro Ala Ile Phe Arg Glu Ala Val Glu Glu Tyr Ile Lys Glu Met
      130                     135                     140
Lys Arg Met Ser Ser Lys Phe Leu Glu Met Val Glu Glu Glu Leu Lys
      145                     150                     155                     160
Ile Glu Pro Lys Glu Lys Leu Ser Arg Leu Val Lys Val Lys Glu Ser
      165                     170                     175
Asp Ser Cys Leu Arg Met Asn His Tyr Pro Glu Lys Glu Glu Thr Pro
      180                     185                     190
Val Lys Glu Glu Ile Gly Phe Gly Glu His Thr Asp Pro Gln Leu Ile
      195                     200                     205
Ser Leu Leu Arg Ser Asn Asp Thr Glu Gly Leu Gln Ile Cys Val Lys
      210                     215                     220
Asp Gly Thr Trp Val Asp Val Thr Pro Asp His Ser Ser Phe Phe Val
      225                     230                     235                     240
Leu Val Gly Asp Thr Leu Gln Val Met Thr Asn Gly Arg Phe Lys Ser
      245                     250                     255
Val Lys His Arg Val Val Thr Asn Thr Lys Arg Ser Arg Ile Ser Met
      260                     265                     270
Ile Tyr Phe Ala Gly Pro Pro Leu Ser Glu Lys Ile Ala Pro Leu Ser
      275                     280                     285
Cys Leu Val Pro Lys Gln Asp Asp Cys Leu Tyr Asn Glu Phe Thr Trp
      290                     295                     300
Ser Gln Tyr Lys Leu Ser Ala Tyr Lys Thr Lys Leu Gly Asp Tyr Arg
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<210> 11

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 11
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<210> 12
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 12
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<210> 13
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<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer

<400> 13
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<210> 14
<211> 20
<212> DNA
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<223> Description of Artificial Sequence: Primer

<400> 14
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<210> 15
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

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<210> 16
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<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer

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